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Sequence Listing could not be accepted.

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Reviewer: Anne Corrigan

Timestamp: [year=2009; month=5; day=27; hr=8; min=47; sec=31; ms=793; ]

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Reviewer Comments:

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<211> 1310

<212> PRT

<213> Varicella zoster

<220>

<221> misc\_feature

<222> (99)..(99)

<223> The 'Xaa' at location 99 stands for Thr, or Met.

<220>

<221> misc\_feature

<222> (512)..(512)

<223> The 'Xaa' at location 512 stands for Ala, or Val.

<220>

<221> misc\_feature

<222> (1275)..(1275)

<223> The 'Xaa' at location 1275 stands for Ser, Leu, or Xaa.

The above <223> response describing "Xaa" at location 1275 contains an error: "Xaa" can only represent an actual amino acid; "Xaa" cannot represent an Xaa.

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Application No: 10591787 Version No: 1.0

Input Set:

Output Set:

Started: 2009-05-19 13:25:27.500  
Finished: 2009-05-19 13:25:45.311  
Elapsed: 0 hr(s) 0 min(s) 17 sec(s) 811 ms  
Total Warnings: 5  
Total Errors: 0  
No. of SeqIDs Defined: 91  
Actual SeqID Count: 91

Error code	Error Description
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W 213	Artificial or Unknown found in <213> in SEQ ID (3)
W 213	Artificial or Unknown found in <213> in SEQ ID (4)
W 213	Artificial or Unknown found in <213> in SEQ ID (7)

# SEQUENCE LISTING

<110> Nagaike, Kazuhiro  
Mori, Yasuko  
Gomi, Yasuyuki  
Takahashi, Michiaki  
Kouichi, Yamanishi

<120> Recombinant varicella-zoster virus

<130> 59150-8037

<140> 10591787

<141> 2009-05-19

<150> PCT/JP2005/003652

<151> 2005-03-05

<150> JP 2004-063277

<151> 2004-03-05

<160> 91

<170> PatentIn version 3.2

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Leu Glu Leu Met Asp Leu Leu Asp Ala Ala Ala Ala Ala Glu His  
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Ser Val Ser Gly Leu Gln Pro Glu Pro Arg Thr Glu Asp Val Gly Glu  
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100 105 110 115	
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Val Ser Leu Gly Thr Lys Leu Ala Arg Ser Pro Lys Pro Pro Met Asn	
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Asp Gly Glu Thr Gly Arg Gly Thr Thr Pro Pro Phe Pro Gln Ala Phe	
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Ile Ser Gly Lys Lys Pro Gly Asp Glu Gln Ala Gly His Ala His Ala	
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Ser Gly Asp Gly Val Val Leu Gln Lys Thr Gln Arg Pro Ala Gln Gly	
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Lys Ser Pro Lys Lys Lys Thr Leu Lys Val Lys Val Pro Leu Pro Ala	
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Arg Lys Pro Gly Gly Pro Val Pro Gly Pro Val Glu Gln Leu Tyr His	
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Val Leu Ser Asp Ser Val Pro Ala Lys Gly Ala Lys Ala Asp Leu Pro	
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Arg Glu His Ala Pro Ser Pro Ser Asn Ser Asp Asp Ser Asp Ser Asn	
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Ala Ala Arg Ala Arg Tyr Glu Ala Ser Thr Glu Pro Xaa Pro Leu Tyr	
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Val Pro Glu Leu Gly Asp Pro Ala Arg Gln Tyr Arg Ala Leu Ile Asn	
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Phe Pro Thr Ala Trp Arg Pro Ala Leu Ser Phe Asp Pro Gly Ala Leu			
790	795	800	
gcg gaa atc gcc gct cgg cgt ccg ggc gga gga gac cga cgg ttt ggt			2685
Ala Glu Ile Ala Ala Arg Arg Pro Gly Gly Gly Asp Arg Arg Phe Gly			
805	810	815	
cca ccc agc gga gtg gag gcg ctg cga cgg agg tgc gcc tgg atg cgg			2733
Pro Pro Ser Gly Val Glu Ala Leu Arg Arg Arg Cys Ala Trp Met Arg			
820	825	830	835
cag atc cca gac ccg gag gat gtg agg ctt ctg atc atc tac gat ccg			2781
Gln Ile Pro Asp Pro Glu Asp Val Arg Leu Leu Ile Ile Tyr Asp Pro			
840	845	850	
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Leu Pro Gly Glu Asp Ile Asn Gly Pro Leu Glu Ser Thr Leu Ala Thr			
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Asp Pro Gly Pro Ser Trp Ser Pro Ser Arg Gly Gly Leu Ser Val Val			
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Gly Val Leu Leu Leu Ser Thr Arg Asp Leu Ala Phe Ala Gly Ala Val			
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Ser Gln Tyr His Val Tyr Val Arg Ala Pro Ala Arg Pro Asp Ala Gln			
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Ala Val Val Arg Trp Pro Asp Ser Ala Val Thr Glu Gly Leu Ala Arg			
980	985	990	995



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Arg Ala Gly Pro Lys	Thr Arg Val Pro Leu	Ser Pro Arg Glu Tyr	
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Glu Ala Ala His Ser	His Arg Ala Ala Asn	Arg Trp Gly Leu Gly	
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Ala Ala Gly Pro Glu	Ala Gly Asp Val Pro	Thr Trp Ala Arg Val	
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Ser Ala Asp Glu Ala	Arg Asn Ala Leu Pro	Pro Ile Pro Arg Val	
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Met Trp Pro Pro Gly	Phe Gly Ala Ala Glu	Thr Val Leu Glu Gly	
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Asp Arg Glu His Ala	Leu Glu Pro Asp Asp	Trp Glu Val Gly Cys	
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